

FIG. 2A

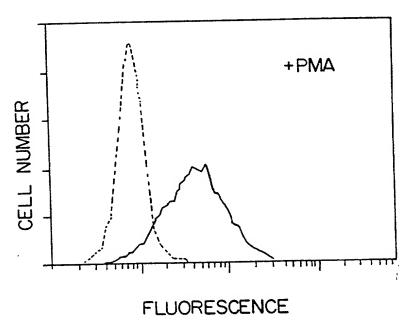


FIG. 2B

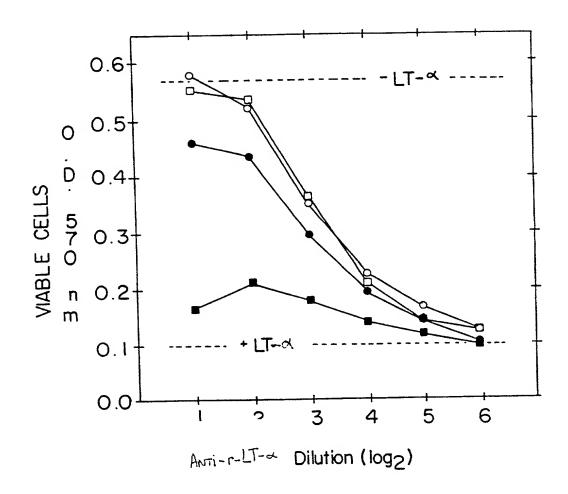


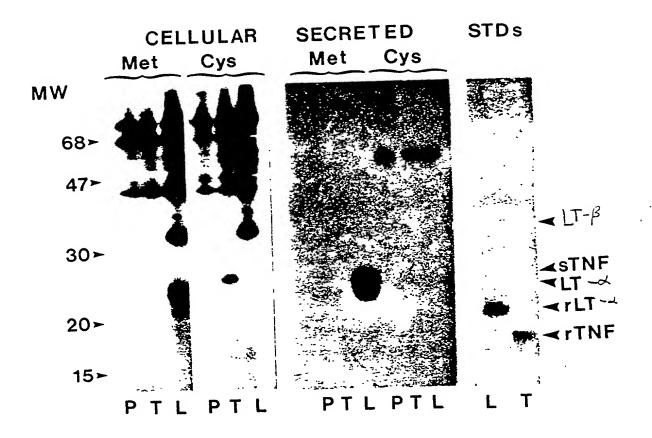
FIG. 3

F1G. 4B

F16. 4A

PRE POST

FIG. 5



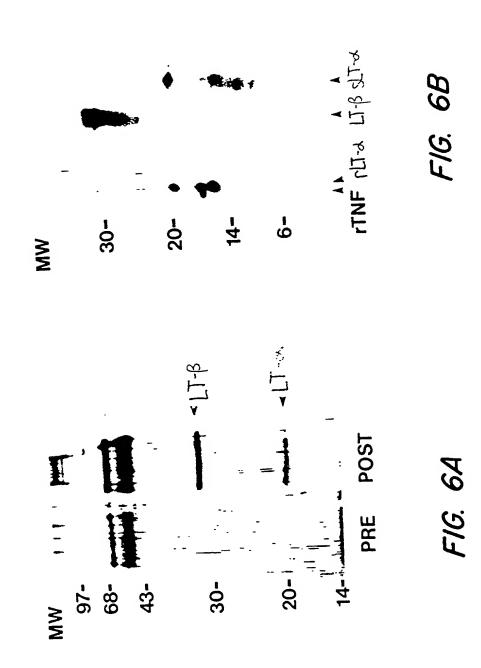




FIG. 7

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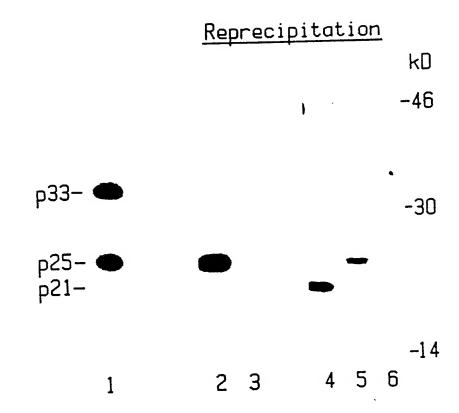


FIG. 8

Denatured

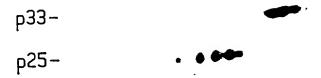


FIG. 9A

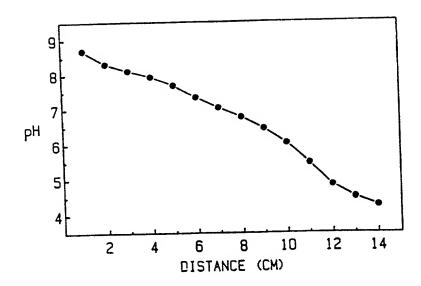


FIG. 9B

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<u>Native</u>

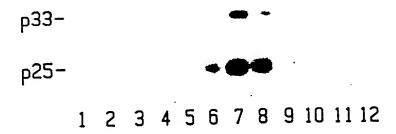


FIG. IOA

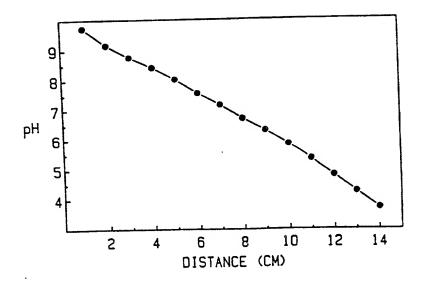
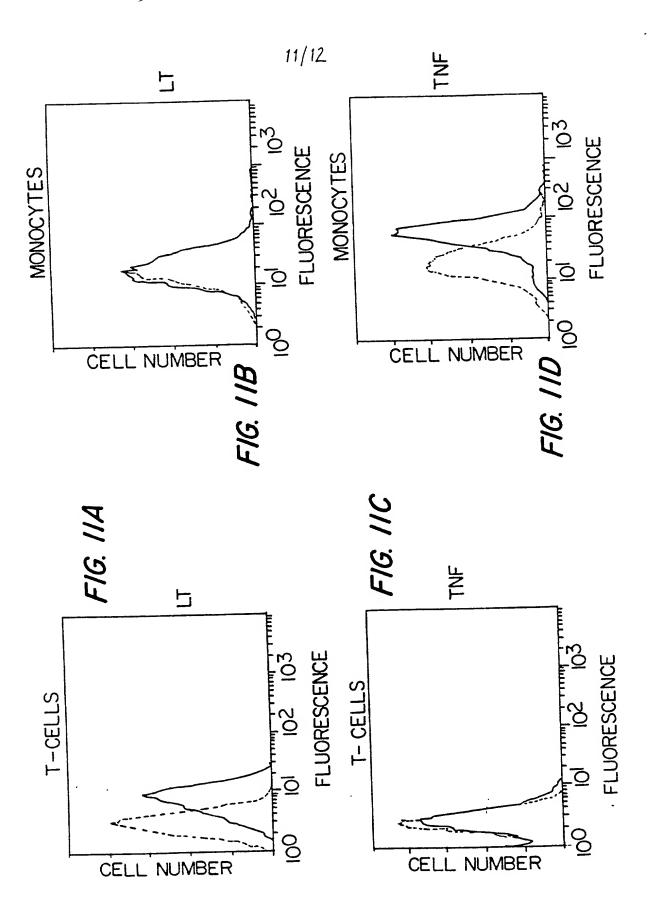
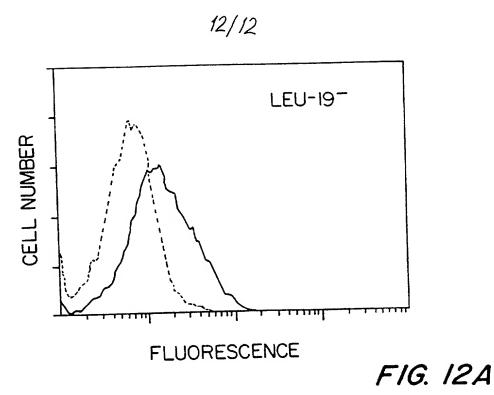
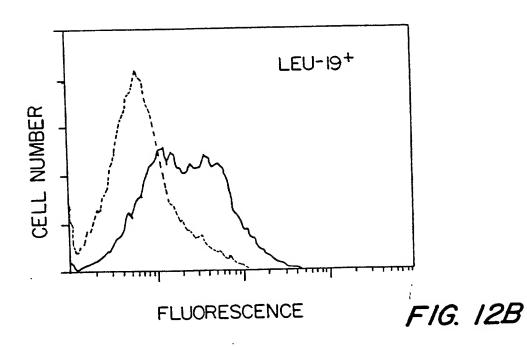


FIG. IOB







LT-β Amino Acid Sequences

N-terminus:

G L E G R G x R L Q G R G S L L L A V A G A T G L V T L L - - V P I

Tryptic Peptides

T87/88:

LPEEEPETDLSPGLPAAHLIGAPLK

x Q A F L T S G T Q F S D A E G L A L P Q D G L Y

XQGLxxET

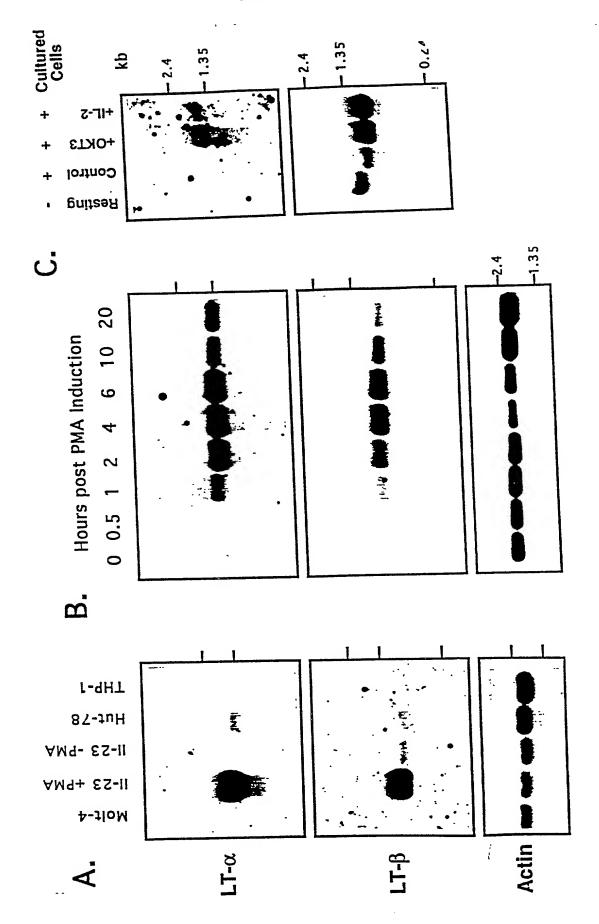
SSLYR

AGGAYGPGTPELLLEGAETVTPVLD

FIGURE 14

SEQUENCE COMPARISON OF VARIOUS LIGANDS IN THE THE-R FAMILY

	1	10	20 	30 1	50 	60
hTNF	MSTESMIRD					
hTNF mCD40L hP33 hLT	MIETYSQP: GLEGRG	SPRSVATGLI GRLQGRGSL	PASMKIFMYL LLAVAGATSL	LTVFLITQMI VTLLLAVPIT	FCLLHFGVIGE GSVLFAVYLHF VLAVLALVPQI PGAQGLPGVGI	RR DQ
hTNF mCD40L hP33 hLT	LDKVEEEVN	LHEDEVEIK				 [T
hTNF mCD40L hP33 hLT	LNKEEKKEN	SFEMQRGDE EEEPETDLS	dpq iaahvvs pgl paahlig	EANSNAASVL APLK-GQ-GL	• QWLNRRANALI QWAKKGYYTM GWETTKEQAFI LWRANTDRAFI	KS L T
hTNF mCD40L hP33 hLT	NLVMLENGK SGTQFSDAE	OLVVPSE GL OLTVKRE GL GLALPOD GL	YYVYTQVTFC YYLYCLVGYR	SNREPSSORP GRAPPGGGDP	* LTHTISRIAVS FIVGLWLKPSI QGRSVTLRSSI SPLYLAHEVQI	IG LY
hTNF mCD40L hP33 hLT	SER RAGGAYGPG	ILLKAAN TPELLLEGA	THSSSQLCEQ ETVTPVLDPA	Q RRQGYGPLWY	•**** EPIYL GGVFQI -SVHL GGVFE I TSVGF GGLVQ I HSMYH GAAFQ I	Le Lo Lr
hTNF mCD40L hP33 hLT	AGASVFVNV RGERVYVNI	TEASQVI SHPD-MVDF	* ••* AESGQVY FGI HRVGFSS FGL AR-GKTF FGA LSPSTVF FGA	L KL V MVG		



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